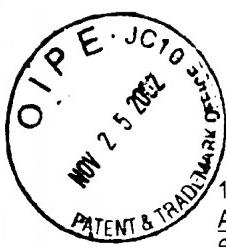


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Figure 1



1 ATGGCGGCGC CGAGCGGACC ATCACCTGGG ACTGGGAGGT TGGCGTCGGT TTACAGCGAG
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241 GGGAGGCAG TGAAGGTCGG GGTGGTGCTC TCTGGTGGAC AAGCACCCGG TGGCACAAT
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361 TTCAAAGGAG GCCCAGCAGG GGTGATGAAG TGCAAGTACG TCAAACCTCAA TACCGATTTC
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541 CTTGTTGTTA TTGGACGGGA CGATTCAAAT ACTCATGCTT GCCTCTTGC TGAATACTTC
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661 AAATGCAAAG AGGTTCCAAC CAGTTTGGA TTTGACACTG CATGCAAGAT CTATTAGAA
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781 CTTATGGGGC GTGCTGCTTC TCACATTACA TTGGGATGCG CTTGCAAAC ACACCCAAAT
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901 TACATTACTG ATATCATCTG CGAGCGTGCA GATCTTGGTT ACAACTATGG TGTTATCCTT
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Figure 2

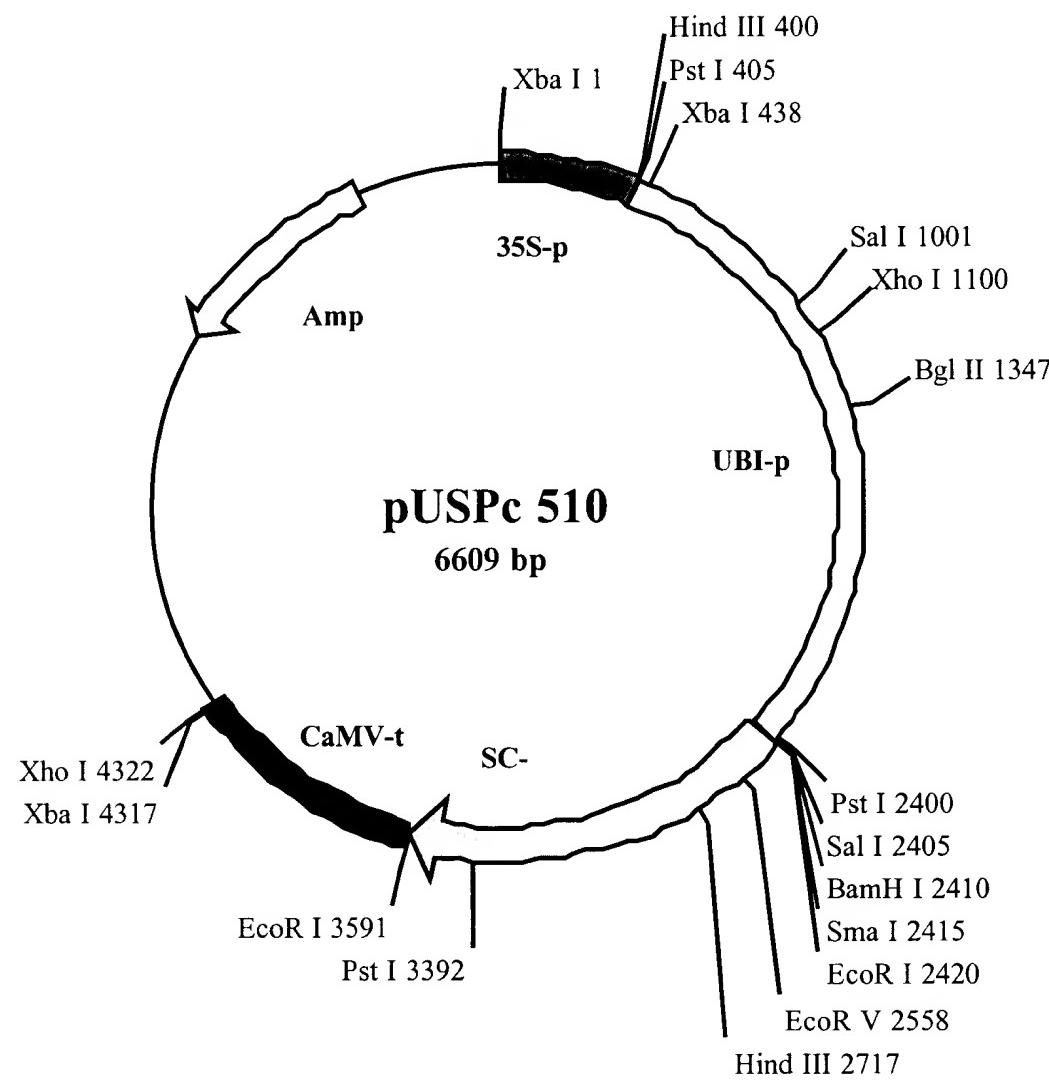


Figure 3

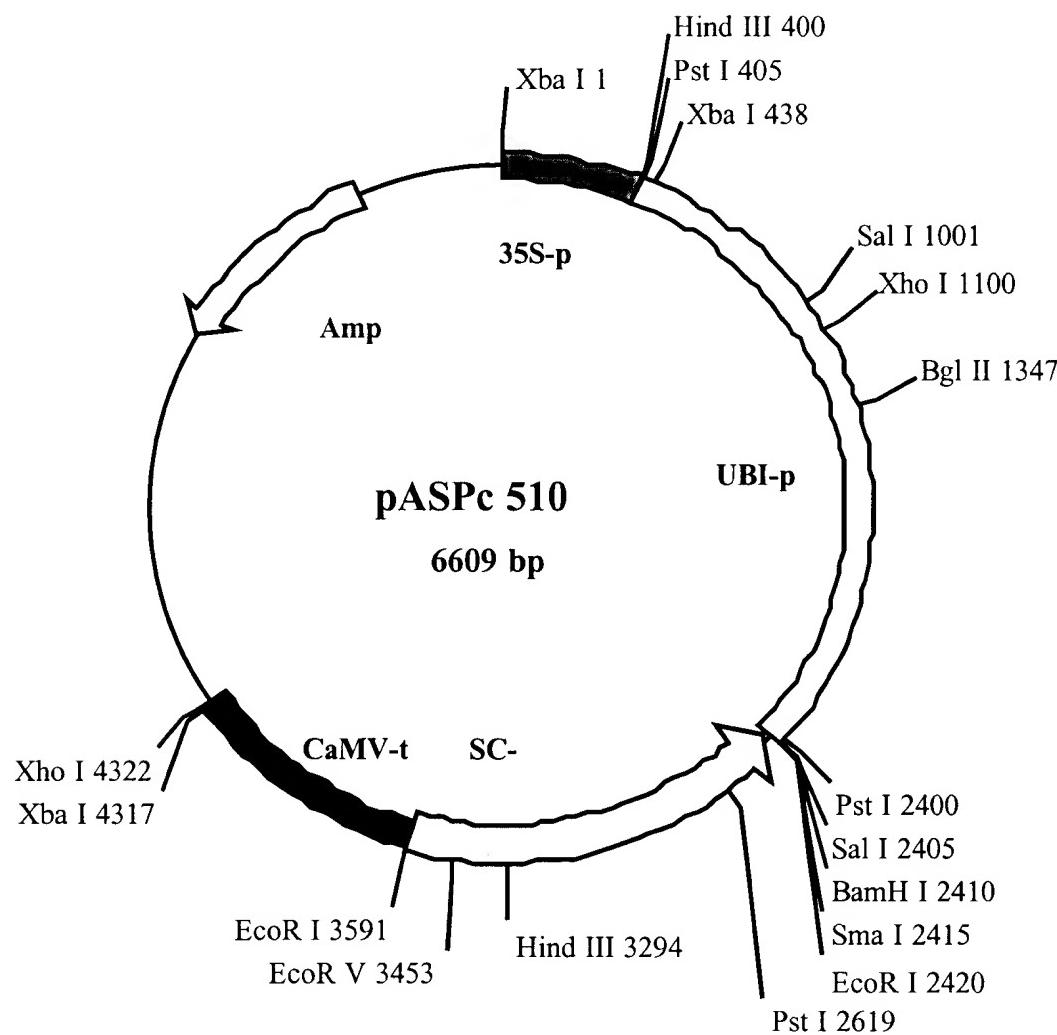


Figure 4



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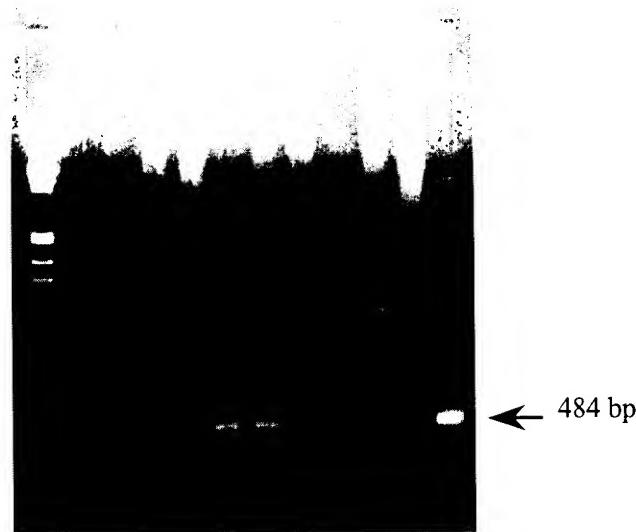


Figure 5

1 2 3 4 5

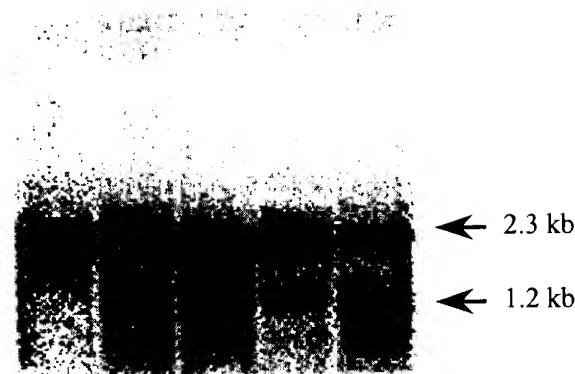


Figure 6

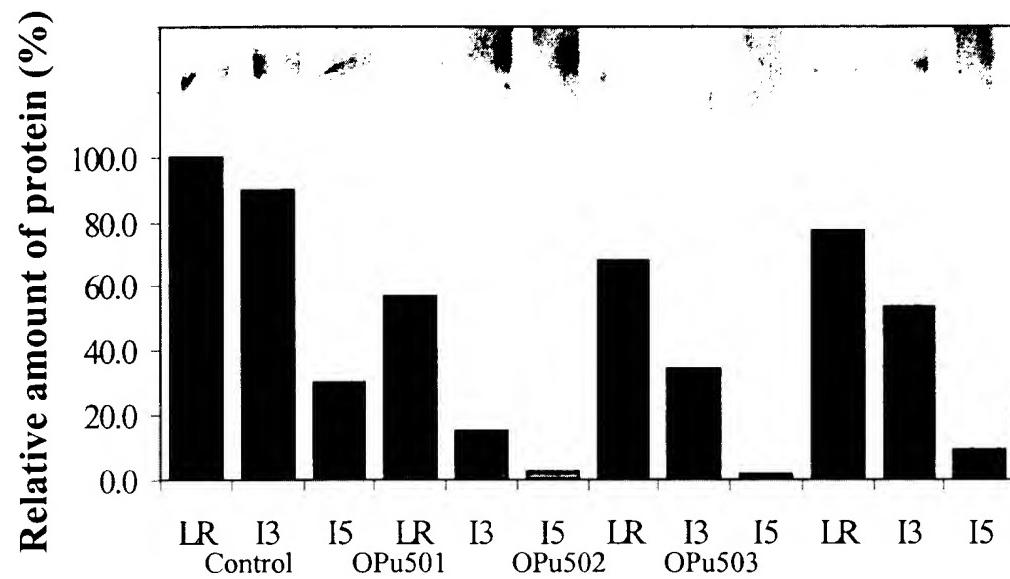
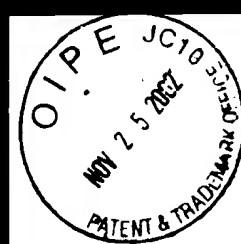


Figure 7